SEQUENCE LISTING

- (1) GENERAL INFORMATION
 - (i) APPLICANT: BRUGE D. GAYNOR, BETTY A. DIAMOND, MATTHEW D. SCHARFF AND PHILIPPE VALADON
 - (ii) TITLE OF INVENTION: PEPTIDES FOR THE TREATMENT AND DIAGNOSIS OF SYSTEMIC LUPUS ERYTHEMATOSUS
 - (iii) NUMBER OF SEQUENCES:
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: AMSTER, ROTHSTEIN & EBENSTEIN
 - (B) STREET: 90 PARK AVENUE
 - (C) CITY: NEW YORK
 - (D) STATE: NEW YORK
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 10016
 - (v) COMPUTER READABLE FORN:
 - (A) MEDIUM TYPE: 3.5 INCH 1.44 Mb STORAGE DISKETTE
 - (B) COMPUTER: IBM PC COMPATIBLE
 - (C) OPERATING SYSTEM: MS-DOS
 - (D) SOFTWARE: ASCII
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/5\\$1,832
 - (B) FILING DATE: SEPTEMBER 20,\1995
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: CRAIG J. ARNOLD
 - (B) REGISTRATION NUMBER: 34,287
 - (C) REFERENCE/DOCKET NUMBER: 96700/342
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (212) 697-5995
 - (B) TELEFAX: (212) 286-0854 or 286-0082
 - (C) TELEX: TWX 710-581-4766
- (2) INFORMATION FOR SEQ ID NO: 1
 - (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 10
- (B) TYPE: AMINO ACID
- (C) STRANDEDNESS: SINGLE
- (D) TOPOLOGY: LINEAR
- (ii) MOLEQULE TYPE:
 - (A) DESCRIPTION: PEPTIDE
- (iii) HYPOTHETICAL: YES
- (iv) ANTI-SENSE: NO
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1

Asp Trp Glu Tyr Ser Val Trp Leu Ser Asn 1 5 10

(3) INFORMATION FOR SEQ ID NO: 2

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (C) STRANDEDNESS: SINGLE
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE:
 - (A) DESCRIPTION: PEPTIDE
- (iii) HYPOTHETICAL: YES
- (iv) ANTI-SENSE: NO
- (ix) FEATURE:
 - (A) NAME/KEY:
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD:
 - (D) OTHER INFORMATION Xaa at 1 and 4 is any amino acid
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2

Xaa Gly Trp Xaa Arg Val

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113886.1



- (4) INFORMATION FOR SEQ ID NO: 3
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
 - (ii) MOLECULE TYPE:
 - (A) DESCRIPTION: PEPTIDE
 - (iii) HYPOTHETIÇAL: YES
 - (ix) FEATURE:
 - (A) NAME/KEY
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD:
 - (D) OTHER INFORMATION: Xaa at 1, 3 and 6 is any amino acid
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3

Xaa Trp Xaa Tyr His Xaa 5

(5) INFORMATION FOR SEQ IN NO: 4

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE:
 - (A) DESCRIPTION: PEPTIDE
- (iii) HYPOTHETICAL: YES
- (ix) FEATURE:
 - (A) NAME/KEY:
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD:
 - (D) OTHER INFORMATION: Xaa at 1 and 3 is Aspartic Acid or Glutamic Acid; Xaa at 5 is Olycine or Serine

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4

Xaa Trp Xaa Tyr Xaa

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- (6) INFORMATION FOR SEQ ID NO: 5
 - (i) SEQUENÇE CHARACTERISTICS:
 - (A) LENGTH: 6
 - (B) TYPE:\ AMINO ACID
 - (D) TOPOLOGY: LINEAR
 - (ii) MOLECULE TYPE:
 - (A) DESCRIPTION: PEPTIDE
 - (iii) HYPOTHETICAL: YES
 - (ix) FEATURE:
 - (A) NAME/KEY:
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD:
 - (D) OTHER INFORMATION: Xaa at 1 and 3 is Aspartic Acid or Glutamic Acid
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5

Xaa Gly Xaa Trp Pro Arg

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- (7) INFORMATION FOR SEQ\ID NO: 6
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
 - (ii) MOLECULE TYPE:
 - (A) DESCRIPTION: PEPTIDE
 - (iii) HYPOTHETICAL: YES
 - (ix) FEATURE:

113886.1

- (A) NAME/KEY:
- (B) LOCATION:
- (C) IDENTIFICATION METHOD:
- (D) OTHER INFORMATION: Xaa at 7-16 is any amino acid
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6

Gly Ala Pro Ser Gly Ala Glu Thr Val

(8) INFORMATION FOR SEQ ID NO: 7

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1d
 - (B) TYPE: AMINO ACID
 - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE:
 - (A) DESCRIPTION: PEPTIDE
- (iii) HYPOTHETICAL: YES
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7

Arg His Glu Asp Gly Asp Trp Pro\Arg Val

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